SEQUENCE LISTING

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<120> METHOD FOR DETERMINING THE SUSCEPTIBILITY OF A SUBJECT TO INFECTION

<130> 3612.1001-000

<150> PCT/EP2003/011240
2003-10-10

<160> 4

<170> PatentIn version 3.0

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             Met Lys Thr Gln Arg Asn Gly His Ser Leu Gly Arg Trp
             1               5                   10

tca ctg gtg ctc ctg ctg ctg ggc ctg gtg atg cct ctg gcc atc att      98
Ser Leu Val Leu Leu Leu Leu Gly Leu Val Met Pro Leu Ala Ile Ile
    15                  20                  25

gcc cag gtc ctc agc tac aag gaa gct gtc ctt cgt gct ata gat ggc     146
Ala Gln Val Leu Ser Tyr Lys Glu Ala Val Leu Arg Ala Ile Asp Gly
30                  35                  40                  45

atc aac cag cgg tcc tcg gat gct aac ctc tac cgc ctc ctg gac ctg     194
Ile Asn Gln Arg Ser Ser Asp Ala Asn Leu Tyr Arg Leu Leu Asp Leu
                50                  55                  60

gac ccc agg ccc acg atg gat ggg gac cca gac acg cca aag cct gtg     242
Asp Pro Arg Pro Thr Met Asp Gly Asp Pro Asp Thr Pro Lys Pro Val
            65                  70                  75

agc ttc aca gtg aag gag aca gtg tgc ccc agg acg aca cag cag tca     290
Ser Phe Thr Val Lys Glu Thr Val Cys Pro Arg Thr Thr Gln Gln Ser
        80                  85                  90

cca gag gat tgt gac ttc aag aag gac ggg ctg gtg aag cgg tgt atg     338
Pro Glu Asp Cys Asp Phe Lys Lys Asp Gly Leu Val Lys Arg Cys Met
    95                  100                 105

ggg aca gtg acc ctc aac cag gcc agg ggc tcc ttt gac atc agt tgt     386
Gly Thr Val Thr Leu Asn Gln Ala Arg Gly Ser Phe Asp Ile Ser Cys
110                 115                 120                 125

gat aag gat aac aag aga ttt gcc ctg ctg ggt gat ttc ttc cgg aaa     434
Asp Lys Asp Asn Lys Arg Phe Ala Leu Leu Gly Asp Phe Phe Arg Lys
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130 135 140

tct aaa gag aag att ggc aaa gag ttt aaa aga att gtc cag aga atc 482
Ser Lys Glu Lys Ile Gly Lys Glu Phe Lys Arg Ile Val Gln Arg Ile
145 150 155

aag gat ttt ttg cgg aat ctt gta ccc agg aca gag tcc tag tgt gtg 530
Lys Asp Phe Leu Arg Asn Leu Val Pro Arg Thr Glu Ser
160 165 170

ccc tac cct ggc tca ggc ttc tgg gct ctg aga aat aaa cta tga gag 578

caa ttt caa aaa aaa aaa aaa aaa aaa acc gga att c 615

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Leu Ser Tyr Lys Glu Ala Val Leu Arg Ala Ile Asp Gly Ile Asn Gln
35 40 45

Arg Ser Ser Asp Ala Asn Leu Tyr Arg Leu Leu Asp Leu Asp Pro Arg
50 55 60

Pro Thr Met Asp Gly Asp Pro Asp Thr Pro Lys Pro Val Ser Phe Thr
65 70 75 80

Val Lys Glu Thr Val Cys Pro Arg Thr Thr Gln Gln Ser Pro Glu Asp
85 90 95

Cys Asp Phe Lys Lys Asp Gly Leu Val Lys Arg Cys Met Gly Thr Val
100 105 110

Thr Leu Asn Gln Ala Arg Gly Ser Phe Asp Ile Ser Cys Asp Lys Asp
115 120 125

Asn Lys Arg Phe Ala Leu Leu Gly Asp Phe Phe Arg Lys Ser Lys Glu
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Lys Ile Gly Lys Glu Phe Lys Arg Ile Val Gln Arg Ile Lys Asp Phe
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Leu Arg Asn Leu Val Pro Arg Thr Glu Ser
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Phe Lys Arg Ile Val Gln Arg Ile Lys Asp Phe Leu Arg Asn Leu Val
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Pro Arg Thr Glu Ser
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